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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULT 1 AF11046 LOCUS DEFINITION ACCESSION VERSION KEYMORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL MEDLINE REFERENCE AUTHORS AUTHORS		C C C C C C C C C C C C C C C C C C C
AF110466 1152 bp N Homo sapiens T-cell cds. AF110466 AF110466.1 GT:43247 human. human. bloaryota; Metazoa; Homo sapiens Eukaryota; Metazoa; 1 (bases 1 to 1152) Pekarsky Y, Hallas, Abnormalities at 14g oncogenes perc. Natl. Acad. Sc 99178995 2 (bases 1 to 1152) Pekarsky Y, Hallas, Sc 99178995 Pekarsky Y, Hallas,		Score Match Length DB ID 1152 100.0 1152 9 AF11 1150.4 99.9 1177 9 AF13 963 83.6 3344 9 AB03 963 83.6 359 9 AB03 718 62.3 114438 2 AC01 701.2 60.9 126308 2 AC01 701.2 60.9 126308 2 AC01 701.2 60.9 126308 9 AF11 699.6 60.7 210791 9 CNS0 699.6 60.7 210791 9 AC01 687.2 59.7 1722 9 AB01 189 16.4 7 753 9 AB02 169.4 14.7 753 9 AB02 62.8 5.5 1240 9 BC00 62.8 5.5 1
mRNA PRI 18-MAR-1999 leukemia/lymphoma 1B (TCLIB) mRNA, complete 04 04 Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo. C., Isobe,M., Russo,G. and Croce,C.M. 32.1 in T cell malignancies involve two i. U.S.A. 96 (6), 2949-2951 (1999) C. and Croce,C.M.	ALIGNMENTS	ID Description

JOURNAL MEDLINE REFERENCE AUTHORS	REFERENCE AUTHORS TITLE	ACCESSION VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 · AF110466 LOCUS DEFINITION
onCogenes Proc. Natl. Acad. Sci. U.S.A. 96 (6), 2949-2951 (1999) 99178995 2 (bases 1 to 1152) Pekarsky,Y., Hallas,C. and Croce,C.M.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1152) Pekarsky,Y., Hallas,C., Isobe,M., Russo,G. and Croce,C.M. Abnormalities at 14q32.1 in T cell malignancies involve two	COS. API10466 API10466.1 GI:4324704 human Homo sapiens	AF110466 1152 bp mRNA PRI 18-MAR-1999 Homo sapiens T-cell leukemia/lymphoma 1B (TCL1B) mRNA, complete

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                                                                                                                                                                                                                                                                                                           Submitted (24-MAR-1999) Biochemistry, Center at Dallas, 5323 Harry Hines Blv
                                                                                                                                                                                                                                                                                                                                                                                                                                      A syncytiotrophoblast-specific gene Syn-1 cloned from human syncytiotrophoblast subtracted cDNA library
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1 (bases 1 to 1177)
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                                                                                                                    901
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     541 gaaggcagtagggcccctgcacactcagtttctctctgttttccttagttatcagtcctgt 600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121
                                                                                                                                                                                                                                         800 GTGCCCAGCCAGCCTCCAGCACCCCCAGTGCAGCTCGTGATTGGAAACTCACCATCGGC 859
                                                                                                                                                                                                                                                                           781 gtgcccagccaggcctccagcacccccagtgcagctcgtgattggaaactcaccatcggc 840
                                                                                                                                                                                                                                                                                                                      740 GCCCCTCATGCTGATCTTCTAGATGCCACTCCCAAATCCCCCTTCATACCCCACCAGGATGT 799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                620 CCTGTCCCACTCAGGTCTGTACTTAGGGCAGCTGGCCTGGATGGGCTTCACTGGGGCCCT 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           481 tgaggatcttcatgtacctgctcttctgtttgcacacccagcatagcctccttgcaggca
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cccgggaggaagtaccgagcagcggattccagtttctgggaaatagcagaccatggccag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ggcagcagatatgaacccagcatcacagtgcacttgtggcagatggcagtgcatacccgg 240
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                                                                                                 atgccctgtgggtatcagttctgctgacactttggcccgaaatagatccagtgctgagca 960
                                                                                                                                                                                 aggcagtggttcggtttaagagatggcattagagggagcccagtctggatgtggacttgg 900
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                                                                                                                                                           AGGCAGTGGTTCGGTTTAAGAGATGGCATTAGAGGGAGCCCAGTCTGGATGTGGACTTGG 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGAGGATCTTCATGTACCTGCTCTTCTGTTTGCACACCCAGCATAGCCTCCTTGCAGGCA 559
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                                                                             ATGCCCTGTGGGTATCAGTTCTGCTGACACTTTGGCCCGAAATAGATCCAGTGCTGAGCA 979
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99.9%;

 Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Saitou,M., Sugimoto,J., Hatakeyama,T., Russo,G. and Isobe,M. Identification of the TCL6 genes within the breakpoint cluster region on chromosome 14932 in T-cell leukemia Oncogene 19 (23), 2796-2802 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AB035343 3144 bp mRNA PRI 12-DEC-2000 Homo sapiens TCL6f4 mRNA for T-cell leukemia/lymphoma 6 ORF105, T-cell leukemia/lymphoma 6 ORF72, complete cds, clone:pDG4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fax:+81-76-445-6874)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T-cell leukemia/lymphoma 6 ORF72; TCL6f4; T-cell leukemia/lymphoma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens cell_line:Daudi cDNA to mRNA, clone:pDG4.
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                                                                                                         AQWGAWRDGFGLCFCLFSPSAHGRDASP"
                                                                                                                            /codon_start=1
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SCHCRFESRPFFLPSPSWSYNRSWGGESAERTWMKQRGTCLSHSPLDLLHPGLCQHKV
                                                                                                                                                                                                                                                                                                                                                                  /codon_start=1
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/protein_id="BAA96383.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                 1805. .2122
/gene="TCL6f4"
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1805. .2122
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/clone="pDG4"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /chromosome="14"
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Score 963;
DB 9;
Length 3144;
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В

Db 3142 AAA 3144 RESULT 4	1150	Oy 1090 tcacagtactggttaattaaacttta	Oy 1030 gtgcttctgcttattcagtccttcag	Oy 970 caccggagcctcagtgagcccatctc	Oy 910 gggtatcagttctgctgacactttgg	Qy 850 ttcggtttaagagatggcattagagg 	Qy 790 caggcctccagcaccccagtgcagc	Qy 730 gctgatcttctagatgccactcccas	Qy 670 ctgagccagtttcccctgctggctgc	Qy 610 ctcaggtctgtacttagggcagctgg 	Qy 550 agggccctgcacactcagtttctct	Oy 490 tcatgtacctgctcttctgtttgcac 	Qy 430 gccctgctggccctgctcttctggc	Qy 370 atggagcagctggtcctaacatatcc 	Oy 310 aagtaccgagcagcggattccagtti	Oy 250 tecteeggceagatgecettetees	Qy 190 tatgaacccagcatcacagtgcact:	Best Local Similarity 100.0%; 1 Matches 963; Conservative 0;
		tcacagtactggttaattaaactttattgctcactgtcaaaaaaaa	gtgcttctgcttattcagtccttcagctcacggaagggatgctagtccgtgaaggtgacc 1089 	caccggagcctcagtgagcccatctgcacagtggggagcatggagggatgggtttggcct 1029 	999tatca9ttct9ct9acacttt99ccc9aaata9atcca9t9ct9a9caa9ca	ttcggtttaagagatggcattagagggagcccagtctggatgtggacttggatgccctgt 909 	caggoctccagcacccccagtgcagctcgtgattggaaactcaccatcggcaggca	gctgatcttctagatgccactcccaaatccccttcatacccaccaggatgtgtgcccagc 789 	ctgagccagtttcccctgctggctgcaagctgtgggttctttct	ctcaggtctgtacttagggcagctggcctggatgggcttcactggggccctgtctgt	agggccctgcacactcagtttctctcgttttcccttagttatcagtcctgtcctgtccca 609 	tcatgtacctgctcttctgtttgcacacccagcatagcctccttgcaggcag	gccctgctggccctgcctcttctggcctggtgtctcctcatgccccctcagtgaggatct 489	atgyagcagctggtcctaacatatcagccggagaggaaagactgacactgggagtggctg 429 	aagtaccgagcagcggattccagtttctgggaaatagcagaccatggccagattgactct 369 	tectocggccagatgccetteteccagetgcccgccgtgtggcagetetaccccgggagg 309	tatgaacccagcatcacagtgcacttgtggcagatggcagtgcatacccgggagctactc 249	<pre>pred. No. 3e-255; ; Mismatches 0; Indels 0; Gaps 0;</pre>
Qy 310 aag Db 2398 AAG	2338	Db 2278 TAT	Matches 96 Oy 190 tat		BASE COUNT		CDS		CDS	9000	Source		AUTHORS TITLE JOURNAL		RS.	SOURCE ORGANISM	Z	AB035342 LOCUS
aagtaccgagcagcggattccagtttctgggaaatagcagaccatggccagattgactct 369 		TATGAACCCAGCATCACAGTGCACTTGTGGGCAGATGGCAGTGCATACCCGGGAGCTACTC 2337 TCCtccggccagatgcccttctcccagctgcccgcgtgtggcagctctacccgggagg 309	3; Conse	83.6%; Score 963; DB 9; Length 3240;	/GL_XEEL="GL: BL/0389" /translation="MALEGAQSGCGLGCPVGISSADTLARNRSSAEQAMYTGASVSPS /AQWGAWRDGFGLCFCLFSPSAHGRDASP" 788 a 833 c 833 g 786 t	/yene-10013 /oodon_start=1 /product="TML1 beta ORE72" /protein_id="BBA96382.1" /http://doi.org/10010101010101010101010101010101010101	SCHERESRPFFLPSSWSYNRSWGGESAERTWMXQRGTCLSHSPLDLLHPGLCQHXV QAR 2951. 3169	/COODI-Start-I /Prodouct="T-cell leukemia/lymphoma 6 ORF105" /protein_id="BAA96381.1" /db_xref="GI:88176588" /db_xref="GI:88176588"	/gene=""CL65" /gene="CL65" /gene="CL65"	/CILLUIGOUNE 14 /Map="14932.1" /Clone="pDG3" /Cell_line="Daudi" 1806 3160	1. 3240 /organism="Homo sapiens" /db_xref="taxon:9606" /organism="1.4"	masanaru isobe, Yuyama university, materiais and Blosystem Engineering, Faculty of Engineering, 3190 Gofuku, Toyama 930-8555, Japan (E-mail:isobe@eng.toyama-u.ac.jp., Tel:+81-76-445-8872, Fax:+81-76-445-6874)	<pre>J (Dases I to 3240) J (Dases I to 3240) ISobe,M., Saitou,M. and Sugimoto,J. Direct Submission Direct Submission Submitted (29-NOV-1999) to the DDBJ/EMBL/GenBank databases.</pre>	region on chromosome 1433 in T-cell leukemia Oncogene 19 (23), 2796-2802 (2000) 20309940	Dukatyold, metazod; Choridatd; Cranidatd; Percebidata; Euteeoscomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 3240) Saitou, M., Sugimoto, J., Hatakeyama, T., Russo, G. and Isobe, M.	Trubels: T-cerl reuxemina/lymphoma o Okrius; Thur beta Okr/2; Thir beta Ok	ll leukemia/lymphoma 6 ORF/2, complete cds, clone:pDG3. 5342. GI:8176587	AB035342 3240 bp mRNA PRI 12-DEC-2000 HOMO Sanions TC16f3 mRNA for T-cell lenkemia/lymphoma 6 OPF105

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Saitou,M.,\ Sugimoto,J.,\ Hatakeyama,T.,\ Russo,G.\ and\ Isobe,M.\ Identification\ of\ the\ TCL6\ genes\ within\ the\ breakpoint\ cluster
                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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                                                                                 Homo sapiens
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                                                                                             Homo sapiens cell_line:Daudi cDNA to mRNA, clone:pDG1.
                                                                                                             leukemia/lymphoma 6 ORF141
                                                                                                                       beta ORF72; TCL6f1; T-cell leukemia/lymphoma 6 ORF72; T-cell
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2930 AGGGCCCTGCACACTCAGTTTCTCTCGTTTTCCTTAGTTATCAGTCCTGTCCTGTCCCA 2989
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                                                                                    2750 ATGGAGCAGCTGGTCCTAACATATCAGCCGGAGAGGAAAGACTGACACTGGGAGTGGCTG 2809
                                                                                                                                                                                                                                                                                                                                          2690 AAGTACCGAGCAGCGGATTCCAGTTTCTGGGAAATAGCAGACCATGGCCAGATTGACTCT 2749
                                                                                                                                                                                                                                                                                                                                                                                                                            2630 TCCTCCGGCCAGATGCCCTTCTCCCAGCTGCCCGCCGTGTGGCAGCTCTACCCCGGGAGG 2689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2570 TATGAACCCAGCATCACAGTGCACTTGTGGCAGATGGCAGTGCATACCCGGGAGCTACTC 2629
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Masaharu Isobe, Toyama University, Materials and Biosystem
Engineering, Faculty of Engineering; 3190 Gofuku, Toyama 930-8555,
Japan (E-mail:isobe@eng.toyama-u.ac.jp., Tel:+81-76-445-6872,
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/gene="TCL6f1"
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/map="14q32.1"
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                                                                                                                                                                                                                                                                       Saltou,M., Sugimoto,J., Hatakeyama,T., Russo,G. and Isobe,M. Identification of the TCL6 genes within the breakpoint cluster region on chromosome 14932 in T-cell leukemia Oncogene 19 (23), 2796-2802 (2000)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                          Fax:+81-76-445-6874)
                                                                                                                                                                               Submitted (29-NOV-1999) to the DDBJ/EMBL/GenBank databases.
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                     /organism="Homo sapiens"
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                                                                      3177 GCTGATCTTCTAGATGCCACTCCCAAATCCCCTTCATACCCACCAGGATGTGTGCCCAGC 3236
                                                                                                                                                            2877 GCCCTGCTGGCCTGCCTTCTGGCCTGGTGTCTCCTCATGCCCCCTCAGTGAGGATCT
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AOWGAWRDGFGLCFCLFSPSAHGRDASP"
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/protein_id="BAA96380.1"
/db_xref="GI:8176586"
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1768. .3528
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/db_xref="GI:8176585"
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1768. .2193
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/map="14q32.1"
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Consensus quality: 105028 bases at least Q40
Consensus quality: 110839 bases at least Q30
Consensus quality: 110839 bases at least Q30
Consensus quality: 111997 bases at least Q20
Estimated insert size: 123000; pulse field gel estimation
Restimated insert size: 113538; sum-of-contigs estimation
Quality coverage: 4.61 in Q20 bases; pulse field gel estimation
Quality coverage: 4.99 in Q20 bases; pulse field gel estimation.
* NOTE: This is a 'working draft' sequence. It currently
* consists of 10 contigs. The true order of the pieces
* is not known and their order in this sequence record is
* arbitrary. Gaps between the contigs are represented as
* runs of N, but the exact sizes of the gaps are unknown.
* This record will be updated with the finished sequence
* as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GTGCTTCTGCTTATTCAGTCCTTCAGCTCACGGAAGGGATGCTAGTCCGTGAAGGTGACC 3536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (15-SEP-1999) Production Sequencing Facility, DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA On May 5, 2000 this sequence version replaced gi:7341483.
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DOE Joint Genome Institute.
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DOE Joint Genome Institute.
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Homo sapiens chromosome 5 clone CTD-2028C21, WORKING DRAFT
SEQUENCE, 10 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                    Summary Statistics
                                                                                                                                                                                                                                                                                                                                                                                           Center clone name: CITB-H1_2028C21
                                                                                                                                                                                                                                                                                                                                                                                                                       Center Project Name: 638370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Center Code: JGI
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                                                                           74534 CCCTCAGTGAGGATCTTTGTGTACTTGCTATTCCGTTTGCACACCCCAGCGTGGCCTCCTT 74475
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                                                                                                     474 ccctcagtgaggatcttcatgtacctgctcttctgtttgccacacccagcatagcctcctt 533
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     534 gcaggcagaaggcagtagggcccctgcacactcagttttctctcgtttttccttagttatca 593
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81.9%; Pred. No. 1.5e-187;
ative 0; Mismatches 155; Indels 54; Gaps
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l14438: contig of 27830 bp in length
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25844: contig of 6088 bp in length
25944: gap of unknown length
37665: contig of 11721 bp in length
37765: gap of unknown length
61723: contig of 23958 bp in length
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                                                                                                                                                                                                                                                                                                                     Direct Submission
Submitted (15-SEP-1999) Production Sequencing Facility, DOE Joint Submitted (15-SEP-1999) Production Sequencing Facility (15-SEP-1999) Production Facility (15-SEP-1999) Product
                                www.jgi.doe.gov
Finishing Completed at Stanford Human Genome Center
www-shgc.stanford.edu
                                                                                                                                                                                                                                                                  DOE Joint Genome Institute and Stanford Human Genome Center
                                                                                                                                                                                                                                                                                               Genome Institute, 2800 Mitchell Drive, Walnut Creek, (bases 1 to 204843)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WI-31186 G24491.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Quality: Phrap Quality >=40 99.7% of Sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               66824 a 40891 c 37481 g 59647 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
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81.9%; Pred. No. 1.5e-187;
ative 0; Mismatches 155; Indels 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .204843
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94947 ATAAAGTGCTGAGCAAGCAGTGTATGCTGGAGCCTCAGTAAGTCCATCTGCACAGTGGGG 94888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1126 tcaaaaaaaaaaa 1139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1006 agcatggagggatgggtttggcctgtgcttctgcttattcagtccttcagctcacggaag 1065
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              atccagtgctgagcaagcaatgtacaccggagcctcagtgagcccatctgcacagtgggg 1005
                                                                                                                                                                                                                                                          Consensus quality: 129558 bases at least Q40 Consensus quality: 135125 bases at least Q30 Consensus quality: 136017 bases at least Q20 Estimated insert size: 143000; pulse field gel estimation Estimated insert size: 137258; sum-of-contigs estimation consists.
* of the gaps between them are based on estimates that have

* provided by the submittor.

* This sequence will be replaced

* by the finished sequence as soon as it is available and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria; Primates; Cat
1 (bases 1 to 137808)
DOE Joint Genome Institute.
Sequencing of Human Chromosome 5
                                                                                                    Estimated insert size: 137258; sum-of-contigs estimation Quality coverage: 7.91 in Q20 bases; pulse field gel estimation Quality coverage: 8.24 in Q20 bases; sum-of-contigs estimation.

* NOTE: This is a 'working draft' sequence. It currently * consists of 12 contigs. Gaps between the contigs.

* are represented as runs of N. The order of the pieces.

* is believed to be correct as given, however the sizes.
                                                                                                                                                                                                                                                                                                                                                                                                         Summary Statistics
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Center clone name: CITB-H1_2221G1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Center: Joint Genome Institute Center Code: JGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (03-AUG-1999) Production Sequencing Facility, DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA On May 6, 2000 this sequence version replaced g1:7709381.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens chromosome 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center Project Name: 712558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Web site: http://www.jgi.doe.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOE Joint Genome Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AC008889.4 GI:7711367
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 137808)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         137808 bp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            clone CTD-2221G1, WORKING DRAFT SEQUENCE
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Db 104859 GA------CTCCTCTCTGGCCTCATGTCTTCGGGC 104887
                                                                                                                                                                                     Db 104803 ----CCAGATCGTCTCCATGGAGCAGCTGGTCCTAACATATTGGCCGGCAAGGAATAACT 104858
                                                                                                                                                                                                                                                                                          Db 104745 AGCTCTACCCCAAGAGGAGGTATCAAGCAGCAAATTCCAGTTTCTGGGAAATAGTGGA-- 104802
                                                                                                                                                                                                                                                                                                                                                                                               Db 104685 ACACCCGGGAACCACTGTCCCCCAGCCAGCTGCCCTTCTCCCCACCTGCTTCCCGTGTGGC 104744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Db 104625 CTCCCAGGGGCACATATGAACCTAGCATCACAGTGCACATGTGGTAGATGCCAGTGC 104684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Db 104565 CTGGGTGACTGTGGTTGTGCGCCCTCAGTCCCCTGCACAGAGAAAGGGCCAGGTGCTCAAG 104624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Db 104505 CCCCCCAACTGTCTGTGGATCTGGAGACCTGGCATCTATGAGGATGACAAGGGGAGGAC 104564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Db 104445 AGGCGACCCCAGAGTGCAGCCTTGCCATGGCCTCCAAAGCTTCTCCGCATCTCAGGGCAC 104504
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BASE COUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 935; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
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                                                                                                                           120 ctgggtgactgtggtcgtgcggttcaatccctcgcgtagggaatgggccagg-----g 172
473 cccctcagtgaggatcttcatgtacctgctcttctgttttgcacacccagcatagcctcct 532
                                                                                                                                                                                                            353 atggccagattgactctatggagcagctggtcctaacatatcagccggagaggaaagact 412
                                                                                                                                                                                                                                                                                                                                    293 agctctaccccgggaggaagtaccgagcagcggattccagtttctgggaaatagcagacc 352
                                                                                                                                                                                                                                                                                                                                                                                                                                         233 atacccgggagctactctcctccggccagatgcccttctcccagctgcccgccgtgtggc 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             173 cctcccagggcagcagatatgaacccagcatcacagtgcacttgtggcagatggcagtgc 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62 cccctgg--ccgtctgtggatccagaggcctggcatctacgaagatgaggaggaggagaac 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 aggegggteeeggttgeagaettgeeatggeeteegaagettetgtgegtetaggggtge 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             * the accession number will be preserved.

1 7204: contig of 7204 bp in length
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116646
118982
119082
132968
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34115
34215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   49909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="CalTech human BAC library D" 25657 c 28235 g 43290 t 1117 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:9606"
/chromosome="5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      133067: gap of unknown length
134154: contig of 1087 bp in length
134254: gap of unknown length
137808: contig of 3554 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 701.6; DB 2;
Pred. No. 5.3e-183;
0; Mismatches 165;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gap of unknown length contig of 13886 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gap of unknown length
contig of 25155 bp in length
gap of unknown length
contig of 2336 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gap of unknown length
contig of 5892 bp in length
gap of unknown length
contig of 9702 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gap of unknown length contig of 2206 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gap of unknown length contig of 24504 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gap of unknown length contig of 21609 bp in length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1117 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels 55; Gaps
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Db 105547 GAAAAAGAAAAAAA 105561
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Db 105368 GATAAAGTGCTGAGC-AGCAGTGTATGCTGGAGCCTCAGTAAGTCCATCTGCACAGTGGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                105487 GGATGCTAATCNCATGAAGGTGACCTCGCAGTACTGGTTAATTAAGCTTTATTGCTCACT 105546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     105248 GGAACTCACCATCAGCAGTCAGTGGCTCGGTTTAAGAGAGGGCCGCAGAGGGGAACTGGGT 105307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1125 gtcaaaaaaaaaaaa 1139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1005 gagcatggagggatgggtttggcctgtgcttctgcttattcagtccttcagctcacggaa 1064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   945 gatccagtgctgagcaagcaatgtacaccggagcctcagtgagcccatctgcacagtggg 1004
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                                                                                                                                              Direct Submission
                                                                                                                                                                           Pekarsky, Y., Hallas, C. and Croce, C.M.
                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 6486)
Pekarsky,Y., Hallas,C., Isobe,M., Russo,G. and Croce,C.M. Abnormalities at 14q32.1 in T cell malignancies involve two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens T-cell leukemia/lymphoma 1B (TCL1B) gene, complete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AF110465.1 GI:4324702
                                                                                                                                                                                                        (bases 1 to 6486)
                                                                                                                                                                                                                                                                      Natl. Acad. Sci. U.S.A. 96 (6), 2949-2951 (1999)
                                                          Location/Qualifiers
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                                                                                       1084 gtgacctcacagtactggttaattaaactttattgctcactgtcaa 1129
                                                                                                                                                                              6228 TGGCCTGTGCTTCTGCTTATTCAGTCCTTCAGCTCACGGAAGGGATGCTAGTCCGTGAAG
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join(141. .329,4439. .4609,4971. .5039,5634. .6330)
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FEATURES

JOURNAL TITLE AUTHORS MEDLINE JOURNAL TITLE

REFERENCE

oncogenes

99178995

REFERENCE

SOURCE KEYWORDS VERSION ACCESSION

ORGANISM

Homo sapiens

DEFINITION AF110465

AF110465

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ch 60.8%; Score 700.4; DB 9; Length 1248; il Similarity 81.5%; Pred. No. 1.2e-182; 928; Conservative 0; Mismatches 156; Indels 54; Gaps 8;	Query Match Best Local Matches 92	
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10th Str. BLSB 1032, Philadelphia, PA 19107, USA		
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roc. Nat1. 1 9178995	JOURNAL MEDLINE	
Pekarsky, r., Hallas, C., ISODe, M., Russo, G. and Croce, C.M. Abnormalities at 14q32.1 in T cell mallgnancies involve two	TITLE	
248)	REFERENCE	
ryota alia;		
human. Homo sapiens	SOURCE	
•	ACCESSION VERSION	
Homo Sapiens T-cell leukemia/lymphoma lB (TCLlB) pseudogene, complete sequence.	DEFINITION	
DDT 18-MAP-100	467	
	RESULT 11	

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1132 AGCGTGGAGGGATGGGTTTGGCTTATGCTTCTGCTTCTGTTCAGCTCTTTCAGCTCACAGCA- 1190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1072 ATAAAGTGCTGAGCAAGCAGTGTATGCTGGAGCCTCAGTAAGTCCATCTGCACAGTGGGG 1131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           952 GAACTCACCATCAGCAGTCAGTGGCTCGGTTTAAGAGAGGGCCGCAGAGGGAACTGGGCT 1011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    826 aaactcaccatcggcagtggttcggtttaagagatggcattagagggagcccagtc 885
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                   1 (bases 1 to 166308)
Heilig,R., Petit,J.L., Vico,V., Dasilva,C., Robert,C., Wincker,P.,
Brottier,P., Cattolico,L., Barbe,V., Pelletier,E., Artiguenave,F.,
Levy,M., Eckenberg,R., Bruls,T., deberardinis,V., Cruaud,C.,
Gyapay,G., Saurin,W. and Weissenbach,J.
Sequencing of the human chromosome 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CNSO1DX2 166308 bp DNA PRI 26-APR-2001
Human chromosome 14 DNA sequence BAC R-164H13 of library RPCI-11
from chromosome 14 of Homo sapiens (Human), complete sequence.
Center: Genoscope / Centre National de Sequencage
                                                                                      Submitted (26-APR-2001) Genoscope - Centre National de Sequencage : BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr
                                                                                                                                     Direct Submission
                                                                                                                                                         Genoscope.
                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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BASE COUNT
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source
                                             y Match 60.7%; Score 699.6; DB 9; Length 166308; Local Similarity 99.4%; Pred. No. 1.9e-182; hes 702; Conservative 0; Mismatches 4; Indels 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The following BAC sequence is oriented from the T7 to the SP6 end Upstream BAC (overlapping the T7 end) : R-1070N10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Percentage of bases with a quality value >= 40 : 99 %.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Downstream BAC (overlapping the SP6 end): R-185P18 (AC=AL133167)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: SeqRef@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center code: GS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        site: http://www.genoscope.cns.fr/
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                                                                                                                                                                                                                             dbSTS:STS26938
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/clone_lib="RPCI-11"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           784 occagocaggoctocagcacococagtgcagoctogtgattggaaactcaccatcggcagg 843
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29130 TGGCCTGTGCTTCTGCTTATTCAGTCCTTCAGCTCACGGAAGGGATGCTAGTCCGTGAAG 29189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28950 CAGTGGTTCGGTTTAAGAGATGGCATTAGAGGGAGCCCAGTCTGGATGTGGACTTGGATG 29009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28830 CCTCATGCTGATCTTCTAGATGCCCACTCCCAAATCCCCCTTCATACCCCACCAGGATGTGTG 28889
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28710 GTCCCACTCAGGTCTGTACTTAGGGCAGCTGGCCTGGATGGGCTTCACTGGGGCCCTGTC 28769
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             964 aatgtacaccggagcctcagtgagcccatctgcacagtggggagcatggagggatgggtt 1023
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                     Submitted (23-MAY-2001) Genoscope - Centre National de Sequencage : BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr
                                                                                                                                                                       Levy,M., Eckenberg,R., Bruls,T., deBerardinis,V., Cruaud,C., Gyapay,G., Saurin,W. and Weissenbach,J. Sequencing of the human chromosome 14
                                                                                                                                                                                                                                        Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 210791)

Hellig,R., Petit,J.L., Vico,V., Dasilva,C., Robert,C., Wincker,P.,
Brottler,P., Cattolico,L., Barbe,V., Pelletter,B., Artiguenave,F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CNS01DV7 210791 bp DNA PRI 23-MAY-2001 Human Chromosome 14 DNA sequence BAC R-1070N10 of library RPCI-11 from chromosome 14 of Homo sapiens (Human), complete sequence.
                                                                         Direct Submission
                                                                                                 Genoscope.
                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Web : www.genoscope.cns.fr)
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STS		STS	STS	STS		STS	STS	STS	cource	FEATURES				COMMENT
EMBL: AA129468	Ω	<pre>dbsTs.STs42889 Identified using the e-PCR software (G. Schuler)" 55639 . 55818 /note="matching EMBL:T72567</pre>	Identified using the e-PCR software (G. Schuler) 755616. 55844 /note="matching EMBL:T72567 RHdb:RH53712	4291. 44412 /note="matching EMBL:AA857911 RHdb:HH93740 dbsTs:STS66733	"matching EMBL: 216446 H95830 H48997 H12943 H7985 STS23702	THOUSE HADD: RAID EMBL: RAID A RHOD: RHOBS1 ABSTS: STS49708 Identified using the e-PCR software (G. Schuler)" 3440034562	· ·	/clone="R-1070N10" /clone_lib="RPCI-11" /139611567 /note="matching EMBL:H04638	/ Organism="Homo sapiens" /db_xref="taxon:9606" /chromosome="14"	90 - 99 : Percentage of Loc	10 - 19 : 10 - 19 : 28567	coverage: 7.83x 1	The following BAC sequence is oriented from the T7 to the SP6 end. Upstream BAC (overlapping the T7 end): C-2588C21 Downstream BAC (overlapping the SP6 end): R-164H13 (AC-AL139020)	On May 30, 2001 this sequence version replaced gi:12657316.
Qy 1024 tggcctgtgcttctgcttattcagtccttcagctcacggaagggatgctagtccgtgaag 1083	Qy 964 aatgtacaccggagcctcagtgagcccatctgcacagtggggagcatggagggatgggtt 1023	QY 904 ccctgtgggtatcagttctgctgacactttggcccgaaatagatccagtgctgagcaagc 963 	Qy 844 cagtggttcggtttaagagatggcattagagggaggcccagtctggatgtggacttggatg 903 		203		Qy 544 ggcagtagggcccttgcacactcagtttctctcgttttccttagttatcagtcctgtcct 603	Qy 484 ggatcttcatgtacctgctcttctgtttgcacacccagcatagcctccttgcaggcag	Qy 424 tggctggccctgctggccctgctcttctggcctggtgtctcctc	Query Match 60.7%; Score 699.6; DB 9; Length 210791; Best Local Similarity 99.4%; Pred. No. 1.9e-182; Matches 702; Conservative 0; Mismatches 4; Indels 0; Gaps 0;	AA856757 PCR software (G. S	Identified using the e-PCR software (G. Schuler)" 189248189446 /note="matching EMBL:R70888 RHdb:RH77821 dbSTS:STS54563 Identified using the e-PCR software (G. Schuler)"	dbsTs:STS38175 Identified using the e-PCR software (G. Schuler)* STS 184387	RHdb:RH48023 dbsTs:STS41082 Identified using the e-PCR software (G. Schuler)" STS 181393181583 /note-"matching EMBL:R68409 RHdb:RH45107

Db 203713 TGGCCTGTGCTTCTGCTTATTCAGTCCTTCAGCTCACGGAAGGGATGCTAGTCCGTGAAG 203772

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                                                                                                                                                                                171 g-----gcctcccagggcagcagatatgaacccagcatcacagtgcacttgtggcagat 224
285 cgtgtggcagctctaccccgggaggaagtaccgagcagcggattccagtttctgggaaat 344
                                                                  196 GCCAGTGCACACCGGGAACCACTGTCCCCCAGCCAGCTGCCCTTCTCCCACCTGCTTCC 255
                                                                                                                                                                                                                                                                        111 ggggagaacctgggtgactgtggtcgtgcggttcaatccctcgcgtagggaatgggccag 170
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DOE Joint Genome Institute and Stanford Human Genome Center.
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                                                                             939 ACAGTGGGGAGCGTGGAGGGATGGGTTTTGGCTTATGCTTCTGCTTGTTCAGTCTTTCAGC 998
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                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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844 cagiggitcggittaagagaiggcattagagggagcccagictggaigitggacitiggaig 903
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 1717)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /product="TCL1 / MTCP1-like 1 (TML1)"
1601. .1606
a 466 c 473 g 438 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="14"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EWARASQGSRYEPSITVHLWQMAVHT_ELLSSGQMPFSQLPAVWQLYPRRKYRAADSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             translation="MASEASVRLGVPPGRLWIQRPGIYEDEEGRTWVTVVVRFNPSRR/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="GI:5478620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product="TCL1 / MTCP1-like 1"
/protein_id="BAA82476.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /gene="TML1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            join(AB025272.1:119. .310,AB025273.1:223. .393,265. .318)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="pCos231"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
1. .1717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oin(AB025272.1:149. .310,AB025273.1:223. .393,265. .318)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.7%; Score 687.6; DB 9; Length 1717; 99.3%; Pred. No. 4.3e-179;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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В

Qy В Qy B δÃ рЬ Qγ

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1580 GTGACCTCACAGTACTGGTTAATTAAACTTTATTGCTCACTGTCCA 1625
               1084 gtgacctcacagtactggttaattaaactttattgctcactgtcaa 1129
                                                         1460 AATGTACACCAGAGCCTCAGTGAGCCCATCTGCACAGTGGGGAGCATGGAGGGATGGGTT 1519
                                                                                                                                                                                                                                                      1340 CAGTGGTTCGGTTTAAGAGATGGCATTAGAGGGAGCCCAGTCTGGATGTGGACTTGGATG 1399
                                                                                                                                           964 aatgtacaccggagcctcagtgagcccatctgcacagtggggagcatggagggatgggtt 1023
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Search completed: November 29, 2001, 08:52:45 Job time: 17206 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Database :
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                      | SIDS2/gcgdata/geneseq/geneseqn/NA1980.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1981.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1981.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1983.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1984.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1985.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1985.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1988.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn/NA1989.DAT:*
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| SIDS2/gcgdata/geneseq/geneseqn/NA199.DAT:*
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Gapop 10.0 , Gapext 1.0
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/SIDS2/gcgdata/geneseg/genesegn/NA2001.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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41.6	41.6	45.6	50	62.8	62.8	62.8	62.8	393.8	701.2	1152	Score	
										100.0	Query Match	dě
										1152	Query Match Length DB	
22	20	21	17	22	17	22	21	22	21	21		
AAD04477	AAV84490	AAA10594	AAT18877	AAI58979	AAT18876	AAI60765	AAA89670	AAH98245	AAA75823	AAA75822	ID	SUMMARIES
Human sphingosine	Human secreted pro	Gene encoding a su	Human TCL-1 gene.	Human polynucleoti	Human TCL-1 cDNA.	Human polynucleoti	Human T cell leuka	Human EST-derived	Nucleotide sequenc	Nucleotide sequenc	Description	S

Croce CM, Pekarsky Y; WPI; 2000-611514/58. P-PSDB; AAB18763.

377 377 377 377 377 377	12 40 14 39 15 39 16 39 17 39 18 39 19 39 19 39 20 38 22 38 22 38 23 38
	24.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0
11097 22 585 20 587 19 1317 22 11778 22 11797 20 4274 15 4325 15 4349 15 4349 15 4349 15 4349 15 4364 15 398 11 1089 21 1089 21 1089 21 1089 21 1089 21 1084 22 1084 22 1085 22 1086 22 108	1512 21 2440 21 2617 21 693 21 693 21 693 22 693 22 693 22 693 22 693 22 496 21 284 21 284 21 284 21 798 22
	1 AAC75966 AAA77441 AAA134812 AAA17669 AAA37065 AAA521493 AAC51493 AAC91565 AAC91565 AAC78292 AAH33714 AAC78292 AAH33714 AAC478292 AAH33714 AAC4794323
P. patens S-adeno Human secreted pr Human secreted pr Human secreted pr Human secreted pr Nucleotide sequen Human secreted pr Nucleotide sequen Human secreted pr Human NMDAR2 rece Human NMDAR2 rece Human NMDAR2 rece Human NMDAR2 rece Human colon cance Sequence encoding A. thaliana LDV11 Human colorectal Mouse plwi gene, Human secreted pr Human secreted pr Cat flea head and	Human ORFX OR Human ORFX OR Human Colon C Human PRO1286 Human PRO1286 Human PRO1286 Human PRO1286 Human PRO1286 Human Inng tu Human lung tu Human cancer Human cancer Human colon C DNA sequence
P. patens S-adenos Human secreted pro Nucleotide sequenc Human secreted pro Nucleotide sequenc Human secreted pro Nucleotide sequenc Human MDAR2 recep Human NMDAR2 recep Human MDAR2 recep Human MDAR2 recep Human MDAR2 recep Human MDAR2 recep Human Delor cancer Sequence encoding A. thaliana LDV115 A. thaliana LDV115 Muman secreted grape, deat flea head and	ORFX ORF1521 ORFX ORF2996 COLON CANCER PRO1286 (UNQ PRO1286 (UNQ COLON SEQUENCE COLON TUNO LUNG TUNO LUNG TUNO LUNG TUNO LUNG TUNO LUNG CANCER LUNG CANCER COLON

ALIGNMENTS

DECITE OF	3	
AAA75822	5822	
ID	AAA75822 stan	AAA75822 standard; DNA; 1152 BP.
XX		
AC	AAA75822;	
X		
ΡŢ	22-JAN-2001 (first entry)	(first entry)
XX		
DE	Nucleotide se	Nucleotide sequence of the human Tc1-1b cDNA.
X		
ΚW	Tcl-1; Tcl-1b	; T cell malignancy; chromosome 14 abnormality; lymphoma;
ΚW	T-cell leukae	T-cell leukaemia; immunodeficiency syndrome; ataxia-teleangiectasia; ss.
X		
S	Homo sapiens.	
X		
FH	Key	Location/Qualifiers
FT	CDS	28414
FT		/*tag= a
FΤ		/product= "Tc1-1b protein"
XX		
PN	WO200055169-A1.	
XX		
PD	21-SEP-2000.	
XX		
ΡF	15-MAR-2000;	15-MAR-2000; 2000WO-US06612.
XX		
PR	15-MAR-1999;	99US-0124714.
X		
PA	(UYJE-) UNIV	(UYJE-) UNIV JEFFERSON THOMAS.
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Matches 1152;
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541 gaaggcagtagggcccctgcacactcagtttctctctgttttccttagtttatcagtcctgt 600
                                                                                                                                                                                                                                                              541 gaaggcagtagggcccctgcacactcagtttctctcgttttccttagttatcagtcctgt 600
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The present sequence encodes a human Tc1-lb protein. The Tc1-lb gene is implicated in the development of T cell malignancies. Fragments o

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                                                                                               Novel nucleic acid of Tcl-1 gene family, Tcl-1b, expressed in low levels in normal bone marrow and peripheral lymphocytes, but activated in T-cell leukaemia and lymphoma, used to identify chromosome 14
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                                                                                    abnormalities
                                                                                                                                                                        WPI; 2000-611514/58.
                                                                                                                                                                                                         Croce CM, Pekarsky Y;
                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tcl-1; Tcl-1b; T cell malignancy; chromosome 14 abnormality; lymphoma;
T-cell leukaemia; immunodeficiency syndrome; ataxia-teleangiectasia; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleotide sequence of the human Tcl-lb gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA75823;
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                                                                                                                                                                                                                                           (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                              15-MAR-1999;
                                                                                                                                                                                                                                                                                                              15-MAR-2000; 2000WO-US06612.
                                                                                                                                                                                                                                                                                                                                                  21-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                  WO200055169-Al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   901 atgccctgtgggtatcagttctgctgacactttggcccgaaatagatccagtgctgagca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      841 aggcagtggttcggtttaagagatggcattagagggagcccagtctggatgtggacttgg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          721 gcccctcatgctgatcttctagatgccactcccaaatccccttcatacccaccaggatgt 780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            721 gcccctcatgctgatcttctagatgccactcccaaatccccttcatacccaccaggatgt 780
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12-OCT-2001
                                                                                                                                                                                                             6288 gtgacctcacagtactggttaattaaactttattgctcactgtcca 6333
                                                                                                                                                                                                                                                                                                                                  1024 tggcctgtgcttctgcttattcagtccttcagctcacggaagggatgctagtccgtgaag 1083
                                                     AAH98245;
                                                                                             AAH98245 standard; cDNA; 697
                                                                                                                                                                                                                                                             1084 gtgacctcacagtactggttaattaaactttattgctcactgtcaa 1129
                                                                                                                                                                                                                                                                                                       6228 tggcctgtgcttctgcttattcagtccttcagctcacggaagggatgctagtccgtgaag
                                                                                                                                                                                                                                                                                                                                                                                                     6168 aatgtacaccggagcctcagtgagcccatctgcacagtggggagcatggagggatgggtt 6227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5988 cccagccaggcctccagcacccccagtgcagctcgtgattggaaactcaccatcggcagg 6047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tcl-1b cDNA sequences are used for detecting a target sequence indicating a chromosome 14 abnormality, such as a (14:14)(q11:q32) translocation or a (14)(q11:q32) inversion. Tcl-1b antisense sequences and antibodies are useful for treating a disease state such as T-cell leukaemia or lymphoma associated with a chromosome 14 abnormality. The Tcl-1b gene and its gene product are useful for treating disease states associated with the Tcl-1b locus on chromosome protein including T-prolymphocytic leukaemias, acute and chronic leukaemias associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6108 ccctgtgggtatcagttctgctgacactttggcccgaaatagatccagtgctgagcaagc 6167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6048 cagtggttcggtttaagagatggcattagagggagcccagtctggatgtggacttggatg 6107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5808 gtcccactcaggtctgtacttagggcagctggcctggatgggcttcactggggccctgtc 5867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5628 tcgcaggccctgctggcctcttctggcctggtgtctcctcatgccccctcagtga 5687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6486 BP; 1600 A; 1559 C; 1687 G; 1639 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                               964 aatgtacaccggagcctcagtgagcccatctgcacagtggggagcatggagggatgggtt 1023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               904 ccctgtgggtatcagttctgctgacactttggcccgaaatagatccagtgctgagcaagc 963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               844 cagtggttcggtttaagagatggcattagagggagcccagtctggatgtggacttggatg 903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               784 cccagccaggcctccagcaccccagtgcagctcgtgattggaaactcaccatcggcagg 843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            664 tgtgtgctgagccagtttcccctgctggctgcaagctgtgggttcttttctcctctgtgcc 723
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the immunodeficiency syndrome ataxia-teleangiectasia (AT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ggcagtagggcccctgcacactcagtttctctcgttttccttagttatcagtcctgtcct 5807
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 697 BP; 135 A; 204 C; 181 G; 175 T; 2 other;
                 616 tctgtacttagggcagctggcctggatgggcttcact-ggggccctgtctgtgtgtggg 674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 246; 1275pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated polypeptide for treatment of diseases, diagnostics, raising antibodies and research use {\boldsymbol{\cdot}}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-JAN-2000; 2000US-0491404.
17-JUL-2000; 2000US-0617746.
03-AUG-2000; 2000US-0631451.
15-SEP-2000; 2000US-0663870.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tomato; monkey; dog; sea urchin; expressed sequence tag; ES diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition; ss.
                                                                                        556 cctgcacactcagtttctctctcgttttccttagttatcagtcctgtcctgtcccgtccagg 615
                                                                                                                                                          Tang YT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JAN-2001; 2001WO-US02687
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264 tctacactgagggcagttggcctggatgggcttcactaggggccccgtctgtgcactggg 323
                                                                     204 cctgcccactcagcttctctcattttcttcacttatcagtcttgtcctgttccactcaaa 263
                                                                                                                                        316 cgagcagcggattccagfttctgggaaatagcagaccatggccagattgactctatggag 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200154477-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; horse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human EST-derived coding sequence SEQ ID NO: 102.
                                                                                                                                                                                                                    94
                                                                                                                                                                                                                                                                                                                                                     l caagcagcaaattccagtttctgggaaatagtgga----ccagatcgtctccatggag 54
                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                   ctggccctgcctcttctggcctggtgtctcctcatgccccctcagtgaggatcttcatgt 495
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                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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A, Zhang J, Werhman T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 393.8; DB Pred. No. 8.4e-96
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 22; Length 697;
                                                                                                                                                                                                                                                                                                                                                                                                                          99; Indels 47; Gaps
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                           Exocytosis pathway protein; Exo protein; antiparkinsonian; antidiabetic; antiallergic; antiasthmatic; noctropic; neuroprotective; anticonvulsant; vulnerary; asthma: inflammation; allergy; Chediak-Higashi syndrome; CHS; alzheimer's disease; Parkinson's disease; Huntington's disease; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA89670 standard; cDNA; 561 BP
                                             New nucleic acids encoding Exo proteins which are useful in the
                                                                       WPI; 2000-482908/42.
                                                                                               Luo X
                                                                                                                     (RIGE-) RIGEL PHARM INC
                                                                                                                                                                                                            26-JAN-1999;
01-FEB-1999;
                                                                                                                                                                                                                                                                                               20-JAN-2000; 2000WO-US01431
                                                                                                                                                                                                                                                                                                                                                 WO200043419-A2.
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                               digestion disorder; wound healing disorder; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human T cell leukaemia/lymphoma l nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA89670;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           564 ggggggat-agntctgctgacactgtggcctgaaatanaaaaagtgctgagcaagcagtg
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                           treatment or prevention of exocytosis-mediated disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                            99US-
                                                                                                                                                                                               99US-0118177.
99US-0118178.
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Disclosure; Page 251; 305pp; English such as asthma, inflammation and allergies

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                                  09-JUL-2000;
19-JUL-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis, treatment or prevention of excytosis mediated disorders such as asthma, inflammation, allergies, Chediak-Higashi Syndrome (CHS), Alzheimer's disease, Parkinson's disease, Huntington's disease, dispersed disorders and wound healing disorders. The nucleic acids, antagonists or agonists of Exo proteins are useful in gene therapy. The nucleic acids are also useful for generating transgenic or knock-out animals which can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence encodes a polypeptide which is associated with the exocytosis pathway. cDNA molecules encoding proteins involved in exocytosis have been isolated by yeast one-hybrid and two-hybrid screening. Novel proteins, termed Exo proteins, have been identified that
                                                                                                                                                                                                                                                                                                                      Human; nootropic; immunosuppressant; cytostatic; gene therapy; cance peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; "Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 561 BP; 110 A; 160 C; 151 G; 135 T; 5 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interact with known exocytosis-associated proteins such as GS27, alpha snap, uncl8-1, vamp3, snap-23, and the rab family of proteins exo proteins and their agonists and antagonists are useful in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAI60765 standard; cDNA; 1235 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                development and screening of therapeutically useful reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Human polynucleotide SEQ ID NO 4754.
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14-SEP-2000;
19-OCT-2000;
                                                                                                              21-JAN-2000;
                                                                                                                                                   26-DEC-2000; 2000WO-US34263
                                                                                                                                                                                        26-JUL-2001.
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                                                                                            25-APR-2000;
                                                                                                                                                                                                                                                                                                         Leukaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            267 gagctgctgccaga 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          328 tccagtttctgggaaatagcagaccatggccagattgactctatggagcagctggtccta 387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             147 ccaagcctgctgcctatcatgtggcagctctaccctgatggacgataccgatcctcagac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                268 ttctcccagctgcccgccgtgtggcagctctaccccgggaggaagtaccgagcggat 327
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Similarity 57.7%;
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2000US-0598042.
2000US-0620312.
2000US-0653450.
2000US-0662191.
2000US-0693036.
                                                                                          2000US-0488725.
2000US-0552317.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAI38642-AAI4213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wang J,
Zhao QA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C.N.S disorders.
                                                         Homo sapiens
                                                                                           TCL-1; chromosome-14; leukaemia; lymphoma; gene therapy; ss.
                                                                                                                                 Human TCL-1 cDNA.
                                                                                                                                                                     23-AUG-1996 (first entry)
                                                                                                                                                                                                         AAT18876;
                                                                                                                                                                                                                                          AAT18876 standard; cDNA; 1324 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1235 BP; 259 A; 359 C; 330 G; 287 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 4754; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P-PSDB; AAM41609
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             such as central nervous system injuries -
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                                                                                                                                                                                                                                                                                                                                                      388 acatatcagccgga 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 268 ttctcccagctgcccgccgtgtggcagctctaccccgggaggaagtaccgagcggat 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     372 gagctgctgccaga 385
                                                                                                                                                                                                                                                                                                                                                                                                        312 tccagtttctggcgcttagtgtaccacatcaagattgacggcgtggaggacatgcttctc 371
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             252 ccaagcctgctgcctatcatgtggcagctctaccctgatggacgataccgatcctcagac 311
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nes 112; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gtgcacttgtggcagatggcagtgcatacccgggagctactctcctccggccagatgccc 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequence data for this patent did not form part of the printed
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Wang Z, Wehrman T, 1
Zhou P, Goodrich R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                     Location/Qualifiers
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57.7%;
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, Yang Y,
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Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer;
                                                       Human polynucleotide SEQ ID NO 1182.
                                                                                           22-OCT-2001 (first entry)
                                                                                                                                 AAI58979;
                                                                                                                                                                   AAI58979 standard; cDNA; 1368 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1324 BP; 293 A; 376 C; 348 G; 307 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New TCL-1 protein and gene associated with chromosome 14 abnormalities - useful to develop prods. for detection, treatment and prevention of diseases such as T-cell leukaemia(s) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-OCT-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A cDNA clone (AAT18876) codes for human TCL-1 protein (AAR94974).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RAGG-) RAGGIO-ITALGENE SPA. (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                    328 tccagtttctgggaaatagcagaccatggccagattgactctatggagcagctggtccta 387
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                                                                                                                                                                                                                                                            370 gagctgctgccaga 383
                                                                                                                                                                                                                                                                                              388 acatatcagccgga 401
                                                                                                                                                                                                                                                                                                                                  310 tccagtttctggcgcttagtgtaccacatcaagattgacggcgtggaggacatgcttctc 369
                                                                                                                                                                                                                                                                                                                                                                                                      250 ccaagcctgctgcctatcatgtggcagctctaccctgatggacgataccgatcctcagac 309
                                                                                                                                                                                                                                                                                                                                                                                                                        268 ttctcccagctgcccgccgtgtggcagctctaccccgggaggaagtaccgagcagcggat 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             208 gtgcacttgtggcagatggcagtgcatacccgggagctactctcctccggccagatgccc 267
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hes 112;
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57.7%;

    Mismatches

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system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1368 BP; 287 A; 395 C; 366 G; 320 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                         specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C.N.S disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to human nucleic acids (AAI57798-AAI61369) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 1182; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             such as central nervous system injuries -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-AUG-2000; 2000US-0653450.
14-SEP-2000; 2000US-0662191.
19-OCT-2000; 2000US-0693036.
29-NOV-2000; 2000US-0727344.
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09-JUL-2000; 2000US-0598042
19-JUL-2000; 2000US-0620312
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HYSE-) HYSEQ INC
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  388 acatatcagccgga 401
                                          367 tccagtttctggcgcttagtgtaccacatcaagattgacggcgtggaggacatgcttctc 426
                                                                             328 tocagtttctgggaaatagcagaccatggccagattgactctatggagcagctggtccta 387
                                                                                                                      307 ccaagcctgctgcctatcatgtggcagctctaccctgatggacgataccgatcctcagac 366
                                                                                                                                                              268 ttctcccagctgccgcgtgtggcagctctaccccgggaggaagtaccgagcagcggat 327
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                                                                                                                                                                                                                                              208 gtgcacttgtggcagatggcagtgcatacccgggagctactctcctccggccagatgccc 267
                                                                                                                                                                                                                                                                                                            Match 5.5%;
Local Similarity 57.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang Z,
Zhou P,
                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Goodrich R,
                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                              Score 62.8;
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                                                                                                                                                                                                                                                                                                                                DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Qian XB,
Yang Y,
                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                              Length 1368;
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                                                                                                                                                                                                                                                                                        0;
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33
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        The human TCL-1 gene (AAT18877) was isolated from the region between 2 sets of breakpoints of approx. 160 kb on the TCL-1 locus of chromosome 14. The TCL-1 gene is expressed at high levels in leukaemic cells carrying a t(14:14)(q11:q32) translocation or an inv(14)(q11:q32) inversion. The gene, probes derived from it, antisense sequences, and the encoded protein (AAR94975) can be used
                                                                                                                                                                                                                        New TCL-1 protein and gene associated with chromosome 14 abnormalities - useful to develop prods. for detection, treatment and prevention of diseases such as T-cell leukaemia(s) and
                                                                                                                                                              Disclosure; Page 69-72; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                      Croce CM, Russo G;
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for the diagnosis or treatment of conditions associated with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-AUG-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT18877;
                                                                                                                                                                                                                                                                                                                                                                                                                 (UYJE-) UNIV JEFFERSON THOMAS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     RAGG-) RAGGIO-ITALGENE SPA
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10134 SRSRTTSTCYTAKYSYSTBSTNCYSDSRGSDNSTAKYASTTTYDAYDARCTTDNSTRCYR 10075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a gene encoding a subunit of the cellulose synthase complex of Vigna angularis. The invention relates to subunits of cellulose synthetic equipment, that can be used to increase the amount of cellulose synthesised by a plant. The proteins and genes encoding them can also be used to improve the properties of the cellulose being
                                                                                                Sequence 10732 BP; 3149 A; 1212 C; 2074 G; 2046 T; 2251 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          produced by a plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; Page 14-21; 32pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A gene encoding a cellulose synthetic equipment – for the improvement in the amount of cellulose synthesised in a plant body % \left( 1\right) =\left\{ 1\right\} =\left\{ 1\right
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vigna angularis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gene encoding a subunit of cellulose synthase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2348 tccagtttctggcgcttagtgtaccaccatcaaggt 2382
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA10594 standard; DNA; 10732 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2288 ccaagcctgctgcctatcatgtggcagctctaccctgatggacgataccgatcctcagac 2347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4922 BP; 1091 A; 1278 C; 1168 G; 1202 T; 183 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MIZU/) MIZUNO K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               increased expression of TCL-1 proteins and/or with chromosome 14 abnormalities, esp. T-cell leukaemia and lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  328 tccagtttctgggaaatagcagaccatggccagat 362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           268 ttctcccagctgcccgccgtgtggcagctctaccccgggaggaagtaccgagcagcggat 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        208 gtgcacttgtggcagatggcagtgcatacccgggagctactctcctccggccagatgccc 267
                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                Conservative 211; Mismatches 238; Indels
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                                                                                                                                                                                                                                                                                         4.0%; Score 45.6; DB 21; 13.2%; Pred. No. 0.047;
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Pred. No. 0.0023
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10074 ASTYDASRRCYSSTYRASRGTBSDDCCTBSRYSCYSYDASRYANCDTBCYTTBAKYRARC 10015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV84490 standard; DNA; 1533 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9714 SSDSTYTBYCSRRCCCYYDAYSCSRYDARCYDACYSYSNSTCYDATBTTSRYSTTYSYYS 9655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9774 TYSSRGYDANSTSRYSSRYSSTYSSDSTYSAKYCAKSTTBTBCYYDAYDACYDAYDANCY 9715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          839 gcaggcagtggttcggtttaagagatggcattagagggagcccagtctggatgtggactt 898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             479 agtgaggatcttcatgtacctgctcttctgtttgcacacccagcatagcctccttgcagg 538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 899 ggatgccctgtgggtatcagttctgctgacactttgg 935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       779 gtgtgcccagccaggcctccagcaccccagtgcagctcgtgattggaaactcaccatcg 838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   719 gtgcccctcatgctgatcttctagatgccactcccaaatccccttcatacccaccaggat 778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             599 gtcctgtcccactcaggtctgtacttagggcagctggactggatgggcttcactggggcc 658
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Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;
06-JUN-1997
                                                                                               06-JUN-1997;
                                                                                                                                                                                                                                                          04-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;
                                                                                                                                                                                                                                                                                                                             10-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted protein gene 80 clone HNFAE54.
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97US-0048899
                               97us-0048896
                                                           97us-0048893
                                                                                        97US-0048884
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97US-0048881
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06-JUN-1997; 06-JUN-1997; 06-JUN-1997; 06-JUN-1997;

97US-0048915. 97US-0048949. 97US-0048964. 97US-0048972. 97us-0049020

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06-JUN-1997;
05-SEP-1997;
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P-PSDB; AAW88613
                                                                                      Shi Y, Soppet DR,
                                                                                                                                               Florence K,
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                               1999-059865/05.
                                                                                                               LA, Carter KC, Dillon PJ, Ebner R, Endress GA
Feng P, Ferrie AM, Fischer CL, Florence C;
Ce K, Greene JM, Hu J, Kyaw H, Lafleur DW;
MOORE PA, Ni J, Olsen HS, Rosen CA, Ruben SM;
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97US-0048897.
97US-0048900.
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97US-0057762.
97US-0057765.
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97US-0048970.
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97US-0048882
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97US-0048883
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97US-0057776
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970S-0057629
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                                                                                      Wei Y,
                                                                                  Young P,
                                                                                      Yu G,
                                                                                         Zeng Z;
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                                                                                                                                                                                                                                                                                                                                                  Cd diagnosed by determining the amount of the new polypeptides in a sample or by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the polynucleotides, based on CC which tissues they are most highly expressed in, and include developing products for the diagnosis or treatment of cancer, neurodegenerative CC disorders, tumours, leukemias, diseases of the immune system, autoimmune CC diseases, hepatic and renal disease, lymphomas, inflammation, allergies, CC restenosis, prostate diseases, obesity, disorders, schizophrenia, CC restenosis, prostate diseases, obesity, disorders involving osteoclasts CC such as osteoporosis, arthritis or mallynancies, diseases of testes, CC lung or thymus, diseaster, and corine disorders, infections and AIDS. The CC polypeptides are also useful for identifying their binding partners. CC The present sequence represents a gene encoding a human secreted protein CC (see descriptor line for gene number and clone identification).
                                                                                                                                                                      Matches
                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to nucleic acid sequences (AAV84411 to AAV84633) encoding human secreted proteins (AAV88534 to AAV88756). The secreted protein gene sequences are deposited with the ATCC under deposit numbers ATCC 97979, 97974, 97975, 97976, 97977, 209007, 209008, 209009, 209010, 209011, 209089, 209081, 209082, 209083, 209085, 209511. Host cells comprising recombinant vectors containing the nucleic acid sequences are used for the recombinant production of the secreted proteins. The polynicleotide and maino acid sequences are useful for are
1045 cagtccttcagctcacggaagggatgctagtccgtgaaggtgacctcacagtactggtta 1104
                                                1354 gaggctatgctttggggggacaggccagaatgaagtcctgggtcaggagcccagctggct 1413
                                                                                                                                                                                                                                                                                                        Sequence 1533 BP; 272 A; 454 C; 495 G; 302 T; 10 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Pathological conditions can be also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 343; 772pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disorders, immune diseases, inflammation or blood disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated human genes and the secreted polypeptides they encode - useful for diagnosis and treatment of e.g. cancers, neurological
                                                                                                               985
                                                                                                                                                                                              Local Similarity
                                                                                                      gagcccatctgcacagtggggagcatggagggatgggtttggcctgtgcttctgcttatt 1044
                                                                                                                                                                      Conservative
                                                                                                                                                                                           3.6%;
53.0%;
                                                                                                                                                                      0
                                                                                                                                                                                           Score 41.6; DB 20; Length 1533; Pred. No. 0.25;
                                                                                                                                                                      Mismatches
                                                                                                                                                                            79;
                                                                                                                                                                         Indels
                                                                                                                                                                         0
                                                                                                                                                                         Gaps
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AAD04477
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                                                                                                                          atherosclerosis; coronary artery disease; dyslipidaemia; diabetes; autoimmune disease; inflammatory disease; multiple sclerosis; T helper-l related disease; chronic obstructive pulmonary disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sphingosine-1-phosphate; S1P; drug screening; therapy; haemostasis; thrombosis; allergic reaction; proliferative disease; cancer; haematopoietic disorder; leukaemia; cardiovascular disease; stroke;
                                                                                                                                                                                                                                                                                                                                                                                              Human sphingosine kinase type 1 (hsk1) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-JUL-2001 (first entry)
                                                         asthma; myocardial infarction; neurodegenerative disorder; wound healing; embryogenesis; anticoagulant; cerebroprotective;
                                                                                                                                                                                                                                                                                                                                 Human; sphingosine kinase type 1; sk1; chromosome 17q25.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AADO4477 standard; cDNA; 1719
vulnerary; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD04477;
                                neuroprotective; antipsoriatic; antiarthritic; cytostatic; cardiant;
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                                                                    Вþ
                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                 The present sequence is human sphingosine kinase type 1 (hsk1) cDNA. The hisk1 gene is located on chromosome 17q25.2. The sk1 converts the substrate sphingosine to sphingosine-1-phosphate (SIP). The sk1 gene and encoded polypeptide are applicable in screening drug candidates particularly inhibitors for preventing or treating disorders such as haemostasis, thrombosis, allergic reactions, proliferative diseases including cancer, haematopoletic disorders such as leukaemia, cardiovascular diseases such as stroke, atherosclerosis and coronary artery disease, dyslipidaemia, diabetes including type I and type II diabetes, autoimmune and inflammatory diseases such as multiple sclerosis, Thelper-1 related diseases, chronic obstructive pulmonary disease, asthma, myocardial infarction, neurodegenerative disorders, natural would healing processes and emborates.
1045 cagtccttcagctcacggaagggatgctcagtccgtgaaggtgacctcacagtactggtta 1104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human sphingosine kinase type I gene for screening drug candidates partLoularly inhibitors used for preventing or treating e.g. atherosclerosis, thrombosis, asthma and diabetes
                                                              1610 gggcccagctgcctatgtaaggccttctagtttgttttgagacccccaccccacgaacca 1669
                                                                                                                              1550 gaggctatgctttggggggacaggccagaatgaagtcctgggtcaggagcccagctggct 1609
                                                                                                                                                                                                                                                                                     Sequence 1719 BP; 305 A; 529 C; 556 G; 329 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Fig 1; 91pp; English.
                                                                                                                                                                                                                                                                                                                      natural wound healing processes and embryogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB; AAE00924
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-OCT-1999;
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                                                                                                                                                                985 gagcccatctgcacagtggggagcatggagggatgggtttggcctgtgcttctgcttatt 1044
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                                                                                                                                                                                                                         Local
                                                                                                                                                                                                        , 68
                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gosink M,
                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9908-0162307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      shown as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product= "Human sphingosine kinase type 1 (hsk1)"
/note= "CDS is specifically claimed in claim 2 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ozak consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note "Translational initiator ATG is in a partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                     53.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Melendez AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 2"
                                                                                                                                                                                                    0;
                                                                                                                                                                                                  Score 41.6; DB Pred. No. 0.26; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Takacs L;
                                                                                                                                                                                                                                    DB 22; Length 1719;
                                                                                                                                                                                                    79;
                                                                                                                                                                                                    Indels
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AAC75966
                                                                                                                                                                                                                                                                  immunostimulant; cardiant; thrombolytic; coagulant; vasotropic; antidabetic; hypotensive; dermatological; immunosuppressive; antifundation; antibacterial; antiviral; antifungal; antirheumatic; antithyroid; and antianaemic. The sequences can be used for determining the presence of or predisposition to, or preventing or treating pathological conditions associated with an ORFX-associated disorder. The
                                   hypertension, hypothyroidism, cholesterol ester storage, systemic lupus erythematosus, severe combined immunodeficiency (SCID), AIDS, viral, bacterial or fungal infection, malaria, autoimmune disorders, asthma, altergies, aplastic anaemia, burns, wounds, bone and cartilage damage, nocturnal haemoglobinuria, antiinflammatory disease; to enhance coagulation; to inhibit thrombosis; and as a contraceptive.
                                                                                                                                                                     nucleic acids can be used to express ORFX proteins in gene therapy vectors. The proteins and nucleic acids may be used to treat cancers, proliferative disorders, neurodegenerative disorders, osteoarthritis, graft vs host disease, cardiovascular disease, diabetes mellitus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vulnerary; antipsoriatic; antiparkinsonian; mootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; cosquilant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antilinflammatory; antibacterial; antifungal; antirheumatic; antithyroid; antivinal; antibacterial; cancer; proliferative disorder; hypertension; neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound;
                                                                                                                                                                                                                                                                                                                                                                                                            antipsoriatic; antiparkinsonian; nootropic; neuroprotective;
osteopathic; anticonvulsant; antiarthritic; immunosuppressant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, which represent the human ORFX open reading frames 1 to 3161. The ORFX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 5; Page 2268-2269; 5507pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         which represent the human ORFX open reading frames 1 to 3161. The ORFX sequences have activities such as: cytostatic; hepatotropic; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neurodegenerative disorders and cardiovascular disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-APR-1999;
05-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; open reading frame; ORFX; detection; cytostatic; hepatotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-602362/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAR-2000; 2000US-0540763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAR-2000; 2000WO-US08621.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thrombosis; contraceptive; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ORFX ORF1521 polynucleotide sequence SEQ ID NO:3041.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-FEB-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC75966 standard; cDNA; 1512 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           damage; cartllage damage; antlinflammatory disease; coagulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB41757
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99US-0127728
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Sequence 1512 BP; 323 A; 408 C; 488 G; 292 T; 1 other;

Query Match

Match 3.6%; Local Similarity 89.8%;

Score 41; DB 2. Pred. No. 0.35;

DB 21; Length 1512;

0;

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RESULT
AAC7744
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          which represent the human ORRY open reading frames I to 3161. The ORRY sequences have activities such as: cytostatic; hepatotropic; vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; osteopathic; anticonvulsant; antitarthritic; immunosuppressant; immunostimulant; cardiant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antificial antitional; antitiona
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAR-1999;
02-APR-1999;
05-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vulnerary; antipsofiatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coaquiant; sostropic; antidabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; antiviral; antibacterial; antifungal; antireumatic; antithyroid; antianaemic; gene therapy; cancer; proliferative disorder; hypertension; neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel nucleic acids and peptides derived from open reading frame useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease -
pathological conditions associated with an ORFX-associated disorder. The
                                                                                                                                                                                                                                                                                                                            AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397,
                                                                                                                                                                                                                                                                                                                                                                                                Claim 5; Page 5164-5166; 5507pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-MAR-2000; 2000WO-US08621.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200058473-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   thrombosis; contraceptive; ss.
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vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotectiv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human ORFX ORF2996 polynucleotide sequence SEQ ID NO:5991.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               damage; cartilage damage; antiinflammatory disease; coagulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000-602362/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB43232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Leach M;
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99US-0127636.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypertension, hypothyroidism, cholesterol ester storage, systemic lupus erythematosus, severe combined immunodeficiency (SCID), AIDS, viral, bacterial or fungal infection, malaria, autoimmune disorders, asthma, allergies, aplastic anaemia, burns, wounds, bone and cartilage damage, nocturnal haemoglobinuria, antiinflammatory disease; to enhance coagulation; to inhibit thrombosis; and as a contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. additionally, N may be used to produce the colon cancer associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis
                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acids can be used to express ORFX proteins in gene therapy vectors. The proteins and nucleic acids may be used to treat cancers, proliferative disorders, neurodegenerative disorders, osteoarthritis.
and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204
                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 3407-3408; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human colon cancer antigen encoding cDNA SEQ ID NO:1894.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          graft vs host disease, cardiovascular disease, diabetes mellitus
                                                                                                                                                                                                                                                    cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ruben SM, Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200122920-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      colorectal carcinoma;
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                                                                                                                                                                                                                                                                                               AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          P-PSDB; AAG75407
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RESULT 15
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Best Local
                                                                             02-JUN-1999;
23-JUN-1999;
23-JUL-1999;
                                        01-SEP-1999;
08-SEP-1999;
15-SEP-1999;
15-SEP-1999;
                                                                                                                                                                                             01-DEC-1998;
16-DEC-1998;
                                                                                                                                                                                                                                     30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                              cytostatic; gene therapy; vaccine; ss.
                                                                                                                                                                                                                                                                                                                                                                       Human; PRO; promotion; inhibition; angiogenesis; cardiovascularisation; diagnosis; trauma; wound; cancer; atherosclerosis; cardiac hypertrophy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2617 BP; 663 A; 672 C; 685 G; 595 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         missing at time of publication, meaning no sequences are present for SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                    08-MAR
                                                                                                                                                                                                                                                                08-JUN-2000
                                                                                                                                                                                                                                                                                           WO200032221-A2
                                                                                                                                                                                                                                                                                                                                                           angiogenic; proliferative; cardiant; cardiovascular; antiatherosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA77669;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA77669 standard; cDNA; 693 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N.B. Pages 666 to 682 and page 7053 of the sequence listing were
                             13-SEP-1999,
                                                                                                                          14 - MAY -
                                                                                                                                       28-APR-
                                                                                                                                                      12-MAR-
                                                                                                                                                                                           16-DEC
                                                                                                                                                                                                                                                                                                                                                                                                                  Human PRO1286 cDNA sequence SEQ ID NO:198.
                                                                                                                                                                                                                                                                                                                                                                                                                                            07-NOV-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       601 cctgtcccactcaggtctgtacttagggcagctggcctggatgggcttcactggggccct 660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             541 gaaggcagtagggcccctgcacactcagttttctctctcgtttttccttagttatcagtcctgt 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    327 g 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              661 g 661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      267 gcgggccgccggtggcgcgggcttggcgcgagctcggcgggcttgggtcccgtggtgcgct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  148 t-actgtcgccccatccctgcgcgcccagcctgccaagcagcgtgccccggttgcaggcg 206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           207 tcatgcagcgggcgcgacccacgctctgggccgctgcgctgactctgctggtgctgctcc 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            481 tgaggatcttcatgtacctgctcttctgtttgcacacccagcatagcctccttgcaggca 540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88 kagcggctgggcgctgaggatcagccgcttcctgcctggattccacagcttcgcgccgtg 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121; Conservative
                                                                                                                                                                               1999;
                                                                                                                                                                                                                                     99WO-US28313
                                        99WO-US20111.
99WO-US20594.
                                                                    99US-0144758.
99US-0145698.
                                                                                             99US-0141037.
                                                                                                                                                     99WO-US05028
99US-0123957
                                                                                                                                                                                             98WO-US25108
98US-0112850
                                                                                                            99WO-US12252
                                                                                                                                      99US-0131445
                                                                                                                                                                               99US-0115554.
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50.2%; Pre
1;
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                                        1113 ttattgctcactgtcaaaaaaaaaaaaaaaaaaaaaaa 1152
                                                                                                                     angiogenic disorders in mammals (e.g. atherosclerosis, cancers and cardiac hypertrophy). For example, the nucleic acids (NCs) and vectors containing them and the PRO polypeptide may be used to treat disorders associated with decreased PRO expression. AAA77510 to AAA77721 and AAB24438 to AAB24435 represent nucleotide and protein sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes nucleic acids encoding PRO polypeptides useful for preventing, diagnosing and treating diagnosing a cardiovascular, endothelial or angiogenic disorder in mammals by modulating cell proliferation, angiogenesis and cardiovascularisation, and in these processes. The and for identifying agonists and antagonists of these processes. The
                                                                                                                                                                                                                                                                      Sequence 693 BP; 184 A; 179 C; 230 G; 100 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acids and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate PRO expression such as cardiovascular, endothelial or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids encoding PRO polypeptides useful for preventing, diagnosing and treating diagnosing a cardiovascular, endothelial or angiogenic disorders in mammals -
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29-OCT-1999;
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                                                                                567 cgggccagggccgagacgcgagtcggatgtggtgaactgaaagaaccaataaaatcatgt 626
627 tcctccaaaaaaaaaaaaaaaaaaaaaaaaaaaaa 666
                                                                                                                                                                                    Local Similarity
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                                                                                                                                                                  Conservative
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Williams PM,
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62.0%;
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                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                    Score 39.2; DB 21; Length 693; Pred. No. 0.78;
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Search completed: November 29, 2001, 08:59:23 Job time: 17559 sec

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
            338.38
348.38
37.66.88
37.2.26.68
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37.2.26.68
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1: /cgn2_6/ptodata/2,

2: /cgn2_6/ptodata/2,

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6: /cgn2_6/ptodata/2,
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            /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/B_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
     US-08-232-463-14
US-08-330-272-1
PCT-USS-13663-1
US-08-330-272-5
PCT-USS-13663-5
US-08-473-981-5
US-08-473-981-6
US-09-460-984-80
US-09-460-984-80
US-09-460-984-80
US-08-433-010-2
US-08-433-010-2
US-08-481-1130-4
US-08-481-1130-4
US-08-481-1130-2
US-08-483-389-2
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PRINCIPLE OF THE PRINCI	33	
TON NUMBER: US/07/9 LICATION US/08/25 SCHELFLINGER, F. G. INVENTION: RECOMBINA SEQUENCES: 52 IDENCE ADDRESS: 52 IBMO Diagonal Road, VA SEQUENCES: 52 IBMO Diagonal Road, VA TYPE: FOLEY & Lardner 1800 Diagonal Road, VA SEQUENCES: 52 IBM PC Compatible RE SYSTEM: PC-DOS/MS E: PATENTION RATA: TION NUMBER: US/08/2 DATE: TION NUMBER: US/07/9 DATE: 101 NUMBER: US/07/9 DATE: 26-AUG-1991 TION NUMBER: 29,768 ACTION INFORMATION: BENT, Stephen A. ATION NUMBER: 29,768 CE/DOCKET NUMBER: 30,0 NICATION INFORMATION: NE: (703)683-419 B99149 FOR SED ID NO: 14: 718 base pairs SOURCE: CHRACTERISTICS: 718 STUDENCE: STAGLE STUDENCE: STAGLE	3289 322 4843 570 1357 1394 1394 1398 3385 3385 902 902 902 2684	289
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                                                                          TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: TCL-1 Gene and Protein a TITLE OF INVENTION: Methods and Compositions
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                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                     CLASSIFICATION: 530 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                             LENGTH: 1324 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                          TELEFAX:
                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                        NAME: Misrock, S. Leslie REGISTRATION NUMBER: 18,
                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                  TOPOLOGY:
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                                                                                                                          (212) 790-9090
(212) 790-8864/9741
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                  FEATURE:
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                                                                                                              MOLECULE TYPE:
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LOCATION:
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                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                         NAME: Misrock, S. Leslie
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Sequence 1, Application PC/TUS9513663 GENERAL INFORMATION:
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CURRENT APPLICATION DATA:
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                                                                                              REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 67
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        310 TCCAGTTTCTGGCGCGTTAGTGTACCACATCAAGATTGACGGCGTGGAGGACATGCTTCTC 369
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                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Paia----
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                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US95/13663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1155 Avenue of the Americas
                                                : (212) 790-9090
(212) 790-8864/9741
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ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 67
TELECOMMUNICATION INFORMATION:
                                                                   2228 GTGCTCTTGCGTCGGGAAGACGTCGTCCTGGGGAGGCCTATGACCCCCACCNAGATAGGC 2287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
  268 ttctcccagctgcccgccgtgtggcagctctaccccgggaggaagtaccgagcagcggat 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                  208 gtgcacttgtggcagattgccattacccgggagctactctcctccggccagattgccc 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    328 tccagfttctgggaaatagcagaccatggccagattgactctatggagcagctggtccta 387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       268 ttctcccagctgcccgccgtgtggcagctctaccccgggaggaagtaccgagcagcgggat 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             208 gtgcacttgtggcagatggcagtgcatacccggggagctactctcctccggccagatgccc 267
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                                                                                                                                                        . Match 4.3%; Score 50; DB 2; Length 4922; Local Similarity 57.4%; Pred. No. 0.0013; less 89; Conservative 0; Mismatches 66; Indels
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                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/330,272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             112; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                         66141 PENNIE
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (212) 790-9090
(212) 790-8864/9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Methods and Compositions
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                                                                                                                                                     0; Gaps
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                                                               US-08-473-981A-5
; Sequence 5, Application US/08473981A
; Patent No. 5629162
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                                              GENERAL INFORMATION:
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TELEX: 66141 PE
INFORMATION FOR SEQ
  APPLICANT:
APPLICANT:
                                                                                                                                                                                           2348 TCCAGTTTCTGGCGCTTAGTGTACCACATCAAGGT 2382
                                                                                                                                                                                                                                                                            2288 CCAAGCCTGCTGCCTATCATGTGGCAGCTCTACCCTGATGGACGATACCGATCCTCAGAC 2347
                                                                                                                                                                                                                                                                                                                                                              2228 GTGCTCTTGCGTCGGGAAGACGTCGTCCTGGGGAGGCCTATGACCCCCACCNAGATAGGC 2287
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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REFERENCE/DOCKET NUMBER: 67
TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                      268 ttctcccagctgcccgccgtgtggcagctctaccccggggaggaagtaccgagcagcggat 327
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CITY: New York
STATE: New Yor
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                     deFougerolles, Antonin R
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Springer, Timothy A
                                                                                                                                                                                                                                                                                                                                                                                                                                                4.3%; Score 50; DB 5; Le ilarity 57.4%; Pred. No. 0.00013; Conservative 0; Mismatches 66;
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(212) 790-8864/9741
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US-08-474-087-5
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                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/473,981A
                                                                                                                                                                                                                                                    APPLICANT: de Fougerolles, Antonin R
APPLICANT: Springer, Timothy A
TITLE OF INVENTION: METHODS OF USING INTERCELLULAR ADHESION MOLECULE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1730 СТАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛА 1759
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                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
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LENGTH: 1817 base pairs
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ADDRESSEE: STERNE, I
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                                                                                                                                    STREET: 11.
              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                               COUNTRY:
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STRANDEDNESS: both
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                                                                                                                                                                        1100 NEW YORK AVENUE, N.W. SUITE 600
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(202) 371-26
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                                                                                                                   USA
PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                             STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KESSLER, GOLDSTEIN & FOX, P.L.L.C
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 Mismatches

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US-09-040-984-80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Patent No. 6210883
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SEQUENCE CHARACTERISTICS:
LENGTH: 1817 base pairs
                                                                                              OPERATIG SYSTEM: DOS
OPERATIG SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/09/040,984
APPLICATION NUMBER: US/09/040,984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1730 CTAAAAAAAAAAAAAAAAAAAAAAAAAA 1759
              ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J
REGISTRATION NUMBER: 31,392
                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Reed, Steven G.
APPLICANT: Wang, TongTong
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR DIAGNOSIS
TITLE OF INVENTION: OF LUNG CANCER
NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1123 ctgtcaaaaaaaaaaaaaaaaaaaaaa 1152
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY:
                                                                                                                                                                                                                                                                                                                               ADDRESSEE: SEED and BERRY LLP STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX:
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                                                                           CLASSIFICATION
                                                                                                                                                                                             COMPUTER:
                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                          STATE: WA
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REFERENCE/DOCKET NUMBER: 210121.456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME:
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                                                                                                                                                                                                                                                                                                               Seattle
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                                                                                                                                                                                           IBM Compatible
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US-08-314-615-2
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-436-983-1
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APPLICANT: Gish, Kurt C
APPLICANT: Wilson, Keith E
TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER,
TITLE OF INVENTION: COMPOSITIONS, AND METHODS OF SCREENING FOR COLORECTAL
TITLE OF INVENTION: CANCER MODULATORS
FILE REFERENCE: A-68431/RMS/DAV
CURRENT APPLICATION NUMBER: US/09/436,983
CURRENT FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.1
SEQ ID No.1
TENTONIC: 7017
                                                                                                                                                                    Sequence 2, Application US/08314615 Patent No. 5525487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match 3.3%; Score 37.6; DB 4; Length 2017; Best Local Similarity 54.3%; Pred. No. 0.33; Matches 76; Conservative 0; Mismatches 64; Indels 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                                                                              GENERAL INFORMATION:
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                                                      APPLICANT: Gallatin, W. Michael
APPLICANT: Vazeux, Rosemay
TITLE OF INVENTION: I-CAM Related Protein
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                             1073 agtccgtgaaggtgacctcacagtactggttaattaaactttattgctcactgtcaaaaa 1132
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                                      CORRESPONDENCE ADDRESS:
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LENGTH: 284 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 2:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gallatin, W. michael APPLICANT: Vazeux, Rosemay TITLE OF INVENTION: 1-CAM Related Protein NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1677 AAAGTTGGCGGGGCTTGGCCTGTGCCCTCAGATTCCGCACCAATAAAGCCTTCAAACTCC 1736
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/314,615
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                                                                                                                                                                                                   STREET: Two Fir
                                                                                                                                                          CITY: Chicago
STATE: Illinois
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                                                                                                                                                                                                                                                  ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Bicknell
                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: BOTUN, Michael F. REGISTRATION NUMBER: 25 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Chicago
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                                                                                                                                                                                                                          Two First National Plaza, 20 South Clark
                                                                                                                                       USA
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US-08-314-362-2
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 1781 base pairs
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FILING DATE: 26-MAY-1992
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US
                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Gallatin, W. Michael
APPLICANT: Vazeux, Rosemay
TITLE OF INVENTION: ICAM-Related Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1737 САЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛА 1766
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TELEPHONE: (312)346-5750
TELEFAX: (312)984-9740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                 PRIOR APPLICATION DATA:
                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Bicknell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: No. 5532127and, Greta E. REGISTRATION NUMBER: 35,302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/07/894,061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
APPLICATION NUMBER: US/08/009,266
                                  CLASSIFICATION: 530
                                                          FILING DATE:
                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                           CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 3.2%;
Local Similarity 63.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57; Conservative
                                                                                                                                                                                                                     60603
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                                                                                                                                                                                                                                                          Illinois
                                                                                                                                                                                                                                                                                                 Street
                                                                                                                                                                                                                                                                                                                       Two First National Plaza, 20 South Clark
                                                                                                                                                                                                                                      USA
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                                                                            US/08/433,010
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27866/30918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 37.2; DB 1; Length 1781; Pred. No. 0.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     33;
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RESULT 13
US-08-245-295-4
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US-08-433-010-2
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INFORMATION FOR SEQ ID NO: 2:
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LENGTH: 1781 base pairs
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REFERENCE/DOCKET NUMBER: 31.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312)346-5750
TELEFAX: (312)34-9740
                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 27-UAN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kilgannon, Patrick D.
TITLE OF INVENTION: ICAM-4 Materials and Methods
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
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FILING DATE: 27-JAN-
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                            APPLICATION NUMBER: US UN APPLICATION NUMBER: US UN APPLICATION DATE: 26-MAY-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: sing
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                                                                                                                                                                CLASSIFICATION: 435
                                                                                                                                                                                                      APPLICATION NUMBER: US/08/245,295
                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                          COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Warshall, O'Toole, Gerstein, Murray & Borun STREET: 233 South Wacker Drive, Suite 6300
                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                       MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              Chicago
                                                                                                                                                                                                                                                                                                                                                      60606
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                                                                                                                                                                                                                                                                                 IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                      United States of America
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                                                            US 07/889,724
                                                                                                                        US 07/827,689
  US 07/894,061
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US-08-481-130-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 1781 base pairs
TYPE: nucleic acid
STRANDEDMESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 4:
     FILING DATE: 27-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/889,724
FILING DATE: 26-MAY-1992
                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1677 AAAGTTGGCGGGGGCTTGGCTGTGCCCTCAGATTCCGCACCAATAAAGCCCTTCAAACTCC 1736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1737 САВАВАВАВАВАВАВАВАВАВАВАВАВА 1766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1123 ctgtcaaaaaaaaaaaaaaaaaaaa 1152
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                                                         APPLICATION NUMBER: US 07/827,689 FILING DATE: 27-JAN-1992
                                                                                                                                          APPLICATION NUMBER: US/08/481,130
                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                   STATE:
                                                                                                                                                                                                                                                                                                                  CITY: Chicago
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/102,852 FILING DATE: 05-AUG-1993
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                                                                                                                                                                                                                                                                  60606-6402
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                                                                                                                                                                                                                                                                                                                               233 South Wacker Drive, 6300 Sears Tower
                                                                                                                                                                                                                                                                                United States of America
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  312-474-0448
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                                                                                                                                                                                                                                                                                                                                               Marshall, O'Toole, Gerstein, Murray & Borun
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US-08~656-984A-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08656984A Patent No. 5753502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 4:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 27-JAN-1992
                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gallatin, W. Michael APPLICANT: Kilgannon, Patrick D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1677 AAAGTTGGCGGGGCTTGGCTGTGCCCTCAGATTCCGCACCAATAAAGCCCTTCAAACTCC 1736
                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ICAM-4 Materials and Methods NUMBER OF SEQUENCES: 42
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APPLICATION NUMBER: US 08/245,
FILING DATE: 18-MAY-194
ATTORNEY_AGENT INFORMATION:
NAME: WILLIAMS, JR. JOSEPH A.
REGISTRATION NUMBER: 38,659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: cDNA
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                                                            CLASSIFICATION:
                                                                         APPLICATION NUMBER: US/08/656,984A FILING DATE:
                                                                                                                                                                                                                                                                           CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
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TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    57; Conservative
                                                                                                                                                                                                                                  RY: United States of America 60606-6402
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.2%; Score 37.2; DB 1; Length 1781; 63.3%; Pred. No. 0.4;
                   US 07/827,689
                                                                                                                                                                                                                                                                                                                                                                                                                Patrick D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US 08/245, 295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US 08/102,852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US 08/009,266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US 07/894,061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27866/32713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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; NAME/KEY: CDS; LOCATION: 16..1659
US-08-656-984A-4
                                                                                     TELEFAX: 312-474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 1781 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/009,266
FILLING DATE: 22-JAN-1993
PRIOR APPLICATION NUMBER: US 08/102,852
FILLING DATE: 05-AUG-1993
PRIOR APPLICATION NUMBER: US 08/245,295
FILLING DATE: 18-MAY-1994
PRIOR APPLICATION NUMBER: US 08/245,295
FILLING DATE: 18-MAY-1994
PRIOR APPLICATION NUMBER: US 08/485,604
APPLICATION NUMBER: US 08/485,604
FILLING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: WILLLAMS, JR. JOSEPH A.
REGISTRATION NUMBER: 27866/33321
TELEPHONE: 312-474-6300
TELEPHONE: 312-474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/889,724
FILING DATE: 26-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/894,061
FILING DATE: 05-JUN-1992
                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                        27866/33321
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Query Match 3.2%; Score 37.2; DB 1; Length 1781; Best Local Similarity 63.3%; Pred. No. 0.4; Matches 57; Conservative 0; Mismatches 33; Indels 0; Gaps

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Search completed: November 29, 2001, 08:49:30 Job time: 16986 sec